

Application No.: 09/464528 Docket No.: BB1205 US NA

## In the Claims

## Claims 1-12 (cancelled)

- 12 13. (currently amended) An isolated nucleic acid fragment having constitutive promoter activity selected from the group consisting of:
- a) an isolated nucleic acid fragment comprising the nucleic acid sequence of SEQ ID NO:6 or SEQ ID NO:14 or a subfragment thereof having constitutive promoter activity; and
- b) an isolated nucleic acid fragment which can hybridize with any of the nucleotide sequences set forth in SEQ ID NO:6 or SEQ ID NO:14 under stringent conditions wherein said stringent conditions comprise washing in 0.1xSSC/0.1% SDS at 65°C. CARL AND ZHONSEN: IS THIS OK? CONDITIONS IN EXAMPLE 3 WERE STRINGENT.
- 13 14. (currently amended) A chimeric gene comprising at least one heterologous nucleic acid fragment operably linked to the isolated nucleic acid fragment of claim 1 13.
- $14 \ \underline{15}$ . (currently amended) An expression construct comprising the chimeric gene of claim  $13 \ 14$ .
  - 15 16. (currently amended) A plant comprising the chimeric gene of claim 13 14.
- $16 \underline{17}$ . (currently amended) The plant of claim  $15 \underline{16}$  wherein said plant is a monocot selected from the group consisting of corn, rice, wheat, barley and palm.
- 17-18. (currently amended) The plant of Claim 16 wherein said plant is a dicot selected from the group consisting of *Arabidopsis*, soybean, oilseed *Brassica*, peanut, sunflower, safflower, cotton, tobacco, tomato, potato, and cocoa.
  - 18 19. (currently amended) The plant of claim 17 18 wherein said plant is soybean.
- 19 20. (currently amended) Seed of the plant as in any one of Claims 15, 16, 17 or 18 16, 17, 18, or 19 wherein said seed comprises in its genome the chimeric gene of claim 13 14.
- 20 21. (currently amended) A method of increasing or decreasing the expression of expressing at least one heterologous nucleic acid fragment in a plant cell which comprises:
  - (a) transforming a plant cell with the chimeric gene of Claim 13 14;
- (b) growing at least one fertile mature plant from the transformed plant cell of step (a);
- (c) selecting at least one plant containing a transformed plant cell wherein the expression of the heterologous nucleic acid fragment is increased or decreased which expresses the heterologous nucleic acid fragment.
- $21 \ \underline{22}$ . (currently amended) The method of Claim  $20 \ \underline{21}$  wherein the plant is a monocot selected from the group consisting of corn, rice, wheat, barley and palm.
- 22 23. (currently amended) The method of Claim 21 wherein the plant is a dicot selected from the group consisting of Arabidopsis, soybean, oilseed Brassica, peanut, sunflower, safflower, cotton, tobacco, tomato, potato, and cocoa.

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23 24. (currently amended) The method of Claim 22 23 wherein the plant is soybean.

- 25. (new) A method of decreasing the expression of an endogenous gene in a plant cell which comprises:
  - (a) transforming a plant cell with the chimeric gene of Claim 14;
- (b) growing at least one fertile mature plant from the transformed plant cell of step (a);
- (c) selecting at least one plant containing a transformed plant cell wherein the expression of the endogenous gene is decreased.
- 26. (new) The method of Claim 25 wherein the plant is a monocot selected from the group consisting of corn, rice, wheat, barley and palm.
- 27. (new) The method of Claim 25 wherein the plant is a dicot selected from the group consisting *of Arabidopsis*, soybean, oilseed *Brassica*, peanut, sunflower, safflower, cotton, tobacco, tomato, potato, and cocoa.
  - 28. (new) The method of Claim 27 wherein the plant is soybean.

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